## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT: Frank, Bruce L. Roberts, Peter C. Goodchild, John Craig, J. Charles Mills, John S. Slade, Andrew Roberts, Noel A. Jupp, Raymond
- (ii) TITLE OF INVENTION: OLIGONUCLEOTIDES SPECIFIC FOR HEPATITIS B VIRUS
- (iii) NUMBER OF SEQUENCES: 56
  - (iv) CORRESPONDENCE ADDRESS:
    - (A) ADDRESSEE: Lappin & Kusmer
    - (B) STREET: 200 State Street
    - (C) CITY: Boston
    - (D) STATE: MA
    - (E) COUNTRY: USA
    - (F) ZIP: 02109
    - (v) COMPUTER READABLE FORM:
      - (A) MEDIUM TYPE: Floppy disk
      - (B) COMPUTER: IBM PC compatible
      - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
  - (vi) CURRENT APPLICATION DATA:
    - (A) APPLICATION NUMBER: US
    - (B) FILING DATE:
    - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:

  - (A) NAME: Kerner, Ann-Louise(B) REGISTRATION NUMBER: 33,523
  - (C) REFERENCE/DOCKET NUMBER: HYZ-041
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: 617-330-1300
    - (B) TELEFAX: 617-330-1311
- (2) INFORMATION FOR SEQ ID NO:1:
  - (i) SEQUENCE CHARACTERISTICS:

	(A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GGTGCGCAG	GA CCAATTTATG	20
(2) INFOR	RMATION FOR SEQ ID NO:2:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CATGGTGCT	IG GTGCGCAGA	19
(2) INFOR	RMATION FOR SEQ ID NO:3:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GAAAAAGTT	TG CATGGTGCTG	20

(2) INF	ORMATION FOR SEQ ID NO:4:	
(i	) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii	) MOLECULE TYPE: DNA	
(iii	) HYPOTHETICAL: NO	
(iv	) ANTI-SENSE: YES	
(xi	) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GAGGTGA	AAA AGTTGCATGG	20
(2) INF	ORMATION FOR SEQ ID NO:5:	
(i	) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii	) MOLECULE TYPE: DNA	
(iii	) HYPOTHETICAL: NO	
(iv	) ANTI-SENSE: YES	
(xi	) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
AGGCAGA	GGT GAAAAGTTG	20
(2) INF	ORMATION FOR SEQ ID NO:6:	
(i	) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 12 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii	) MOLECULE TYPE: DNA	
(iii	) HYPOTHETICAL: NO	
(iv	ANTI-SENSE: YES	
(xi	) SEQUENCE DESCRIPTION: SEQ ID NO:6:	

AGGCAGAG	GT GA	12
(2) INFO	RMATION FOR SEQ ID NO:7:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:7:	
AGAGATGA	TT AGGCAGAGGT	20
(2) INFO	RMATION FOR SEQ ID NO:8:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:8:	
GACATGAA	CA AGAGATGATT AGGCAGAGGT	30
(2) INFO	RMATION FOR SEQ ID NO:9:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	
GACATGTA	CA AGAGATGATT	20
(2) INFO	RMATION FOR SEQ ID NO:10:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
GTAGGACA	TG AACAAGAGAT	20
(2) INFO	RMATION FOR SEQ ID NO:11:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	
TTGGAGGC	IT GAACAGTAGG	20
(2) INFO	RMATION FOR SEQ ID NO:12:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	

(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CACAGCTT	GG AGGCTTGAAC	20
(2) INFO	RMATION FOR SEQ ID NO:13:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:13:	
AGCCACCC	AA GGCACAGCTT	20
(2) INFO	RMATION FOR SEQ ID NO:14:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:14:	
TCGATGTC	CA TGCCCCAAAG	20
(2) INFO	RMATION FOR SEQ ID NO:15:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	

(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:15:	
TAAGGGTC	GA TGTCCATGCC	20
(2) INFO	RMATION FOR SEQ ID NO:16:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:16:	
TTATAAGG	GT CGATGTCCAT	20
(2) INFO	RMATION FOR SEQ ID NO:17:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 27 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:17:	
AAATTCTT'	TA TAAGGGTCGA TGTCCAT	27
(2) INFO	RMATION FOR SEQ ID NO:18:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:18:	
TAT	AAGGG'	TC GA	12
(2)	INFO	RMATION FOR SEQ ID NO:19:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:19:	
AAA'	TTCTT	TA TAAGGGTCGA	20
(2)	INFO	RMATION FOR SEQ ID NO:20:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:20:	
GTA!	TCTAG	AA GATCTCGTAC	20
(2)	INFO	RMATION FOR SEQ ID NO:21:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid	

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:21:	
GCGGTGTC'	TA GAAGATCTCG	20
(2) INFO	RMATION FOR SEQ ID NO:22:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:22:	
GAGGCGGT	GT CTAGGAGATC	20
(2) INFO	RMATION FOR SEQ ID NO:23:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:23:	
GAGCTGAG	GC GGTGTCTAGG	20
(2) INFO	RMATION FOR SEQ ID NO:24:	

(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:24:	
ATACAGAG	CT GAGGCGGTAT	20
(2) INFO	RMATION FOR SEQ ID NO:25:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:25:	
TCCCGATA	CA GAGCTGAGGC	20
(2) INFO	RMATION FOR SEQ ID NO:26:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:26:	
AGGCTTCC	CG ATACAGAGCT	20

(2) INFO	RMATION FOR SEQ ID NO:27:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:27:	
ACAATGCT	CA GGAGACTCTA	20
(2) INFO	RMATION FOR SEQ ID NO:28:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:28:	
GCAGTATG	GT GAGGTGAGCA	20
(2) INFO	RMATION FOR SEQ ID NO:29:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:29:	

GAGTGCAG	TA TGGTGAGGTG	20
(2) INFO	RMATION FOR SEQ ID NO:30:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:30:	
TGCCTGAG	TG CAGTATGGTG	20
(2) INFO	RMATION FOR SEQ ID NO:31:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:31:	
TTGCTTGC	CT GAGTGCAGTA	20
(2) INFO	RMATION FOR SEQ ID NO:32:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:32:	
TAAGGGTC	GA AGAGATGATT	20
(2) INFO	RMATION FOR SEQ ID NO:33:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:33:	
AGAGATGA	TT TAAGGGTCGA	20
(2) INFO	RMATION FOR SEQ ID NO:34:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:34:	
TAAGGGTC	GA AGGCAGAGGT	20
(2) INFO	RMATION FOR SEQ ID NO:35:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	

(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:35:	
AGGCAGAG	GT TAAGGGTCGA	20
(2) INFO	RMATION FOR SEQ ID NO:36:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:36:	
TATAAGGG	IC GAAGGCAGAG GTGA	24
(2) INFO	RMATION FOR SEQ ID NO:37:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:37:	
AGGCAGAG	GT GATATAAGGG TCGA	24
(2) INFO	RMATION FOR SEQ ID NO:38:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	

(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:38:	
AGAGATGA	TT AGGCAGAGGT TAAGGGTCGA	30
(2) INFO	RMATION FOR SEQ ID NO:39:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:39:	
GACATGAA	CA AGAGATGATT TAAGGGTCGA	30
(2) INFO	RMATION FOR SEQ ID NO:40:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:40:	
AGAGATGA'	TT TAAGGGTCGA TGTCCATGCC	30
(2) INFO	RMATION FOR SEQ ID NO:41:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:		
AGGCAGAG	GT TAAGGGTCGA TGTCCATGCC	30	
(2) INFO	RMATION FOR SEQ ID NO:42:		
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:		
GGCATTTG	GT GGTCTATAAG	20	
(2) INFO	RMATION FOR SEQ ID NO:43:		
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear		
(ii)	MOLECULE TYPE: DNA		
(iii)	HYPOTHETICAL: NO		
(iv)	ANTI-SENSE: YES		
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:		
GATAGGGG	CA TTTGGTGGTC	20	
(2) INFO	(2) INFORMATION FOR SEQ ID NO:44:		
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs		

	<ul><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:44:	
TGTTGATA	GG ATAGGGGCAT	20
(2) INFO	RMATION FOR SEQ ID NO:45:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:45:	
ACCCAAGG	CA CAGCTTGGAG	20
(2) INFO	RMATION FOR SEQ ID NO:46:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:46:	
GACAGGGG	CA TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:47:	

(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:47:	
GATAGGGG	CC TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:48:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:48:	
GATAGGGG	CA TTTGGTGCTC	20
(2) INFO	RMATION FOR SEQ ID NO:49:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:49:	
GACAGGGG	ግር ጥጥጥርርጥርርጥር	20

(2) INFO	RMATION FOR SEQ ID NO:50:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 20 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:50:	
ATGTTTTT	GG CGTCTTCCAT	20
(2) INFO	RMATION FOR SEQ ID NO:51:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 21 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:51:	
TGCTGGGG	GG AACTAATGAC T	21
(2) INFO	RMATION FOR SEQ ID NO:52:	
(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 25 base pairs  (B) TYPE: nucleic acid  (C) STRANDEDNESS: single  (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: NO	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:52:	

GGA	GTGCG	AA TCCACACTCC GAAAG	25
(2)	INFO	RMATION FOR SEQ ID NO:53:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:53:	
TAA	GGGTC	GA	10
(2)	INFO	RMATION FOR SEQ ID NO:54:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:54:	
AGG	CAGAG	GT	10
(2)	INFO	RMATION FOR SEQ ID NO:55:	
	(i)	SEQUENCE CHARACTERISTICS:  (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:	
ATATTCGCAC GAATTCAGCT TATAGACCAC CAAATG	36
(2) INFORMATION FOR SEQ ID NO:56:	
<ul> <li>(i) SEQUENCE CHARACTERISTICS:</li> <li>(A) LENGTH: 34 base pairs</li> <li>(B) TYPE: nucleic acid</li> <li>(C) STRANDEDNESS: single</li> <li>(D) TOPOLOGY: linear</li> </ul>	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: NO	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:	
GTTGGGATTG CTGCAGCTAT CTGGATTTGC GGTG	34